

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/980,364

CRF Processing Date: 5/31/2002

Edited by: [Signature]

Verified by: [Signature] (STIC staff)

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form. 3/1/95



PCT09

RAW SEQUENCE LISTING

DATE: 06/02/2002

PATENT APPLICATION: US/09/980,364

TIME: 17:15:43

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\05312002\I980364.raw

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3 <110> APPLICANT: Boutilier, Kim
4   Ouellet, Therese
5   Custers, Jan
6   Hattori, Jiro
7   Miki, Brian
8   Van Lookeren Campagne, Michiel
10 <120> TITLE OF INVENTION: Use of the BNM3 Transcriptional Activator to Control
11   Plant Embryogenesis and Regeneration Processes
13 <130> FILE REFERENCE: 270.62USWO
15 <140> CURRENT APPLICATION NUMBER: 09/980,364
C--> 16 <141> CURRENT FILING DATE: 2002-04-08
18 <150> PRIOR APPLICATION NUMBER: EP 99201745.9-2106
19 <151> PRIOR FILING DATE: 1999-06-02
21 <160> NUMBER OF SEQ ID NOS: 14
23 <170> SOFTWARE: PatentIn Ver. 2.1
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 2014
27 <212> TYPE: DNA
28 <213> ORGANISM: Brassica napus
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80 35 40 45
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83 50 55 60
85 Ala Phe Thr Arg Asp Asn Asn Ser His Ser Arg Asp Trp Asp Ile Asn
86 65 70 75 80
88 Gly Cys Ala Cys Asn Asn Ile His Asn Asp Glu Gln Asp Gly Pro Lys
89 85 90 95
91 Leu Glu Asn Phe Leu Gly Arg Thr Thr Thr Ile Tyr Asn Thr Asn Glu
92 100 105 110
94 Asn Val Gly Asp Gly Ser Gly Ser Gly Cys Tyr Gly Gly Gly Asp Gly
95 115 120 125
97 Gly Gly Gly Ser Leu Gly Leu Ser Met Ile Lys Thr Trp Leu Arg Asn
98 130 135 140
100 Gln Pro Val Asp Asn Val Asp Asn Gln Glu Asn Gly Asn Ala Ala Lys
101 145 150 155 160
103 Gly Leu Ser Leu Ser Met Asn Ser Ser Thr Ser Cys Asp Asn Asn Asn
104 165 170 175
106 Asp Ser Asn Asn Asn Val Val Ala Gln Gly Lys Thr Ile Asp Asp Ser
107 180 185 190
109 Val Glu Ala Thr Pro Lys Lys Thr Ile Glu Ser Phe Gly Gln Arg Thr
110 195 200 205
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113 210 215 220
115 Ala His Leu Trp Asp Asn Ser Cys Lys Arg Glu Gly Gln Thr Arg Lys
116 225 230 235 240
118 Gly Arg Gln Val Tyr Leu Gly Gly Tyr Asp Lys Glu Glu Lys Ala Ala
119 245 250 255
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131 305          310          315          320
133 Gln His Gly Arg Trp Gln Ala Arg Ile Gly Arg Val Ala Gly Asn Lys
134          325          330          335
136 Asp Leu Tyr Leu Gly Thr Phe Gly Thr Gln Glu Glu Ala Ala Glu Ala
137          340          345          350
139 Tyr Asp Ile Ala Ala Ile Lys Phe Arg Gly Leu Thr Ala Val Thr Asn
140          355          360          365
142 Phe Asp Met Asn Arg Tyr Asn Val Lys Ala Ile Leu Glu Ser Pro Ser
143          370          375          380
145 Leu Pro Ile Gly Ser Ala Ala Lys Arg Leu Lys Glu Ala Asn Arg Pro
146 385          390          395          400
148 Val Pro Ser Met Met Met Ile Ser Asn Asn Val Ser Glu Ser Glu Asn
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152          420          425          430
154 Asp Leu Ser Leu Leu His Gln His Gln Glu Arg Tyr Asn Gly Tyr Tyr
155          435          440          445
157 Tyr Asn Gly Gly Asn Leu Ser Ser Glu Ser Ala Arg Ala Cys Phe Lys
158          450          455          460
160 Gln Glu Asp Asp Gln His His Phe Leu Ser Asn Thr Gln Ser Leu Met
161 465          470          475          480
163 Thr Asn Ile Asp His Gln Ser Ser Val Ser Asp Asp Ser Val Thr Val
164          485          490          495
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167          500          505          510
169 Val Asn Cys Asp Ala Tyr Ala Ala Ser Glu Phe Asp Tyr Asn Ala Arg
170          515          520          525
172 Asn His Tyr Tyr Phe Ala Gln Gln Gln Gln Thr Gln Gln Ser Pro Gly
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175 Gly Asp Phe Pro Ala Ala Met Thr Asn Asn Val Gly Ser Asn Met Tyr
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190 <400> SEQUENCE: 3
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193 cttccaccac cacaaccgcc gtagatgtcg ccggagagta ctgttacgat ccgaccgctg 180
194 cctccgatga gtcttcagcc atccaaacat cgtttccttc tccctttggt gtcgtcctcg 240

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197 ccaccacgat ttacaacacc aacgaaaacg ttggagatat cgatggaagt ggggtgttatg 420
198 gaggaggaga cgggtgtggt ggctcactag gactttcgat gataaagaca tggctgagaa 480
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200 tctcaatgaa ctcatctact tcttgtgata acaacaacta cagcagtaac aacctgttg 600
201 cccaagggaa gactattgat gatagcgttg aagctacacc gaagaaaact attgagagtt 660
202 ttggacagag gacgtctata taccgcggtg ttacaaggca tcggtggaca ggaagatatg 720
203 aggcacattt atgggataat agttgtaaac gagaaggcca aacgcgcaaa ggaagacaag 780
204 tttatttggg aggttatgac aaagaagaaa aagcagctag ggcttatgat ttagccgcac 840
205 tcaagtattg gggaaaccacc actactacta acttcccat gagcgaatat gagaaagaga 900
206 tagaagagat gaagcacatg acaaggcaag agtatgttgc ctcaattcgc aggaaaagta 960
207 gtggtttctc tcgtgttgca tcgatttatc gtggagtaac aagacatcac caacatggaa 1020
208 gatggcaagc taggatagga agagtcgccc gtaacaaaga cctctacttg ggaacttttg 1080
209 gcacacaaga agaagctgca gaggcatacg acattgcggc catcaaattc agaggattaa 1140
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240 35 40 45
242 Ser Ala Ile Gln Thr Ser Phe Pro Ser Pro Phe Gly Val Val Val Asp
243 50 55 60
245 Ala Phe Thr Arg Asp Asn Asn Ser His Ser Arg Asp Trp Asp Ile Asn
246 65 70 75 80
248 Gly Cys Ala Cys Asn Asn Ile His Asn Asp Glu Gln Asp Gly Pro Lys
249 85 90 95
251 Leu Glu Asn Phe Leu Gly Arg Thr Thr Thr Ile Tyr Asn Thr Asn Glu
252 100 105 110

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261 145      150      155      160
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266 Asp Ser Asn Asn Asn Val Val Ala Gln Gly Lys Thr Ile Asp Asp Ser
267      180      185      190
269 Val Glu Ala Thr Pro Lys Lys Thr Ile Glu Ser Phe Gly Gln Arg Thr
270      195      200      205
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276 225      230      235      240
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279      245      250      255
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282      260      265      270
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285      275      280      285
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291 305      310      315      320
293 Gln His Gly Arg Trp Gln Ala Arg Ile Gly Arg Val Ala Gly Asn Lys
294      325      330      335
296 Asp Leu Tyr Leu Gly Thr Phe Gly Thr Gln Glu Glu Ala Ala Glu Ala
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299 Tyr Asp Ile Ala Ala Ile Lys Phe Arg Gly Leu Thr Ala Val Thr Asn
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303      370      375      380
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306 385      390      395      400
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314 Asp Leu Ser Leu Leu His Gln His Gln Glu Arg Tyr Asn Gly Tyr Tyr
315      435      440      445
317 Tyr Asn Gly Gly Asn Leu Ser Ser Glu Ser Ala Arg Ala Cys Phe Lys
318      450      455      460
320 Gln Glu Asp Asp Gln His His Phe Leu Ser Asn Thr Gln Ser Leu Met
321 465      470      475      480
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/980,364

DATE: 06/02/2002

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Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\05312002\I980364.raw

L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date